

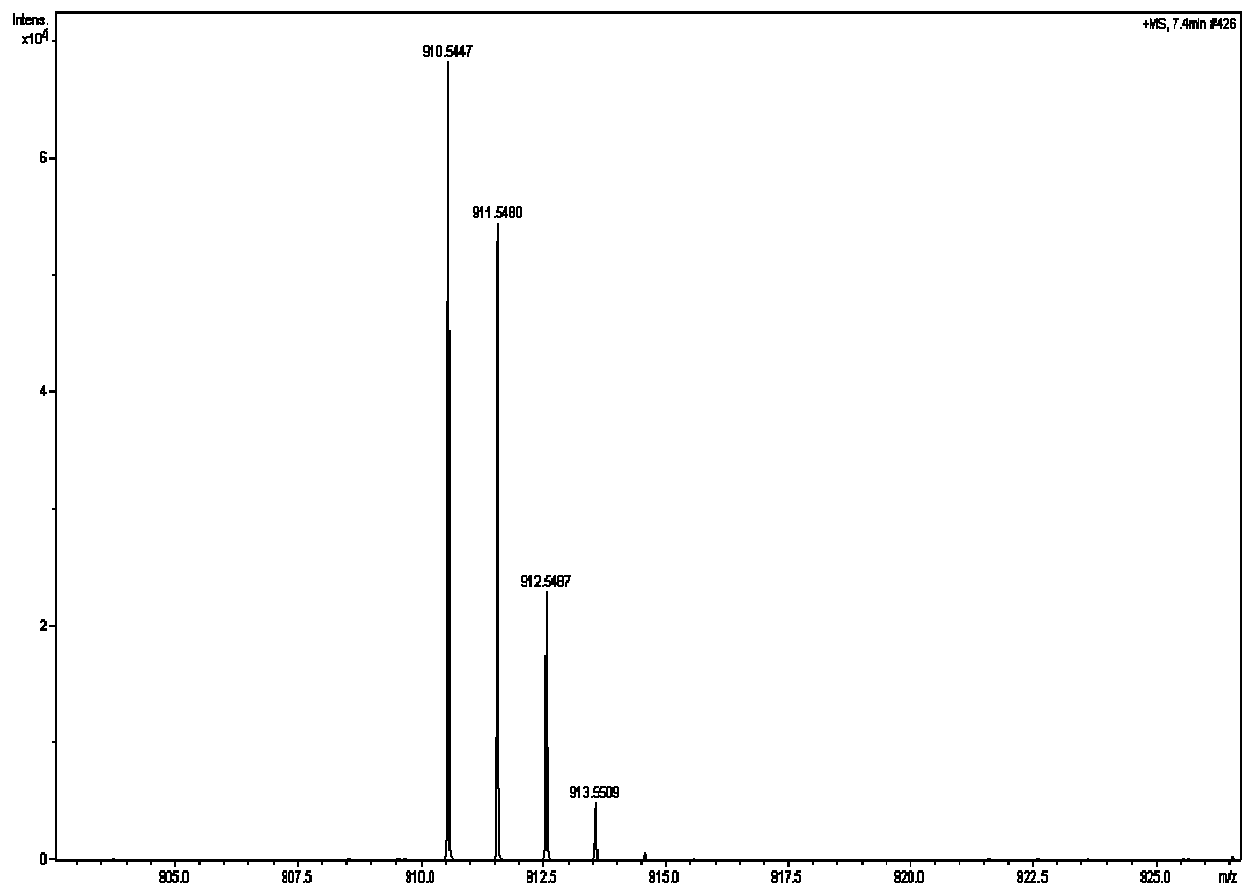
**Hydrogen-deuterium exchange mass spectrometry reveals the interaction of  
Fenna-Matthews-Olson protein and chlorosome CsmA protein**

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**Supporting information**

Mass spectrum of the sample showing a base peak at  $m/z$  401.22. The x-axis represents  $m/z$  from 350 to 450, and the y-axis represents relative intensity. The spectrum shows a single prominent peak at  $m/z$  401.22, which is the base peak.

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**Figure S2.** ESI spectrum of Bchl *a*, which is usually eluted at high organic phase of the HPLC.